

GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

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Run on: February 16, 2003, 15:49:44 ; Search time 215.022 Seconds
        (without alignments)
        14704.597 Million cell updates/sec
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Title: US-09-497-967-3

Perfect score: 1404
Sequence: 1 atgaaaaataatatatttttagt.....tgattttctattattattatta 1404

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 2185239 seqs, 1125999159 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_101002.*

1:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1980.DAT.*
2:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT.*
3:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1982.DAT.*
4:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1983.DAT.*
5:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1984.DAT.*
6:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1985.DAT.*
7:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1986.DAT.*
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9:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1988.DAT.*
10:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1989.DAT.*
11:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1990.DAT.*
12:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1991.DAT.*
13:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1992.DAT.*
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17:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1996.DAT.*
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22:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT.*
23:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT.*
24:	SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	1404	100.0	1404	21	AAA97038	55k i-antigen nuc
2	1404	100.0	1410	21	AAA97060	55k i-antigen cod
3	1402.4	99.9	1404	21	AAA52136	55 kDa i-antigen g
4	782.6	55.7	1404	21	AAA97040	55k i-antigen syn
5	782.6	55.7	1410	21	AAA97089	Synthetic I. Multi
6	781	55.6	1404	21	AAA97005	Synthetic 55k i-a
7	252.6	18.0	1326	21	AAA97036	48k i-antigen nuc
8	252.6	18.0	2486	21	AAA97037	Nucleotide sequenc
9	251	17.9	1326	21	AAA97035	48 kDa i-antigen q

ALIGNMENTS

RESULT 1
AAA97038

ID AAA97038 standard; DNA; 1404 BP.

AC AAA97038;

18-DEC-2000 (first entry)

DE 55kD i-antigen nucleotide sequence.

Immobilisation antigen: i-antigen; ichthyophthiriasis; vaccine; ds;
 KW white spot disease; freshwater fish; immune response; infection control.

OS *Ichthyophthirius multifiliis*.

PN WO200046373-A1.

10-AUG-2000.

04-FEB-2000; 2000WO-US02962.

04 - FEB - 1999; 99US-0118634.

PR 02-MAR-1999; 99US-0122372.
PR 17-MAR-1999; 99US-0124905.

PR 99US-0131121.
27-APR-1999;
XX

PA (UYGE-) UNIV GEORGIA RES FOUND INC.
PA (CORR) CORNELL RES FOUND INC

PA (CLAR/) CLARK T G.
PA (DICK/) DICKERSON T

PA (LINT//) LINT T.
XX

white spot disease; freshwater fish; immune response; infection control.

Ichthyophthirius multifiliis.

W0200046373-A1.

10-AUG-2000.

04-FEB-2000; 2000WO-US02962.

04-FEB-1999; 99US-0118634.

02-MAR-1999; 99US-012372.

17-MAR-1999; 99US-0124905.

27-APR-1999; 99US-0131121.

(UYGE-) UNIV GEORGIA RES FOUND INC.

(CORR.) CORNELL RES FOUND INC.

(CLAR/) CLARK T G.

(DICK/) DICKERSON H W.

(LINT/) LIN T.

Clark TG, Dickerson HW, Lin T;

WPI; 2000-506071/45.

Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius

multifiliis, useful for prophylaxis and treatment of Ichthyophthirius

infection in fish

Disclosure: Figure 2; 144pp: English.

This invention relates to novel i-antigen polypeptide sequences. I-antigens or immobilisation antigens are common to a variety of hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in Ichthyophthirius multifiliis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kD. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis serotypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic i-antigen polypeptide obtained is also useful for treating or preventing I. multifiliis infection in fish. Sequences AAA97036-A97042, and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene fragments identified in the invention. Sequences AAA97043-A97064 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent i-antigen protein and peptide sequences.

Sequence 1410 BP; 449 A; 240 C; 259 G; 462 T; 0 other;

Query Match 100.0%; Score 1404; DB 21; Length 1410;
Best Local Similarity 100.0%; Pred. No. 1.5e-301;
Matches 1404; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ATGAAAAATATATTTAGTAAATTTGATTTATTTTCAATTTATCAATTAATTAATCT 60
Db 1 ATGAAAAATATATTTAGTAAATTTGATTTATTTTCAATTTATCAATTAATTAATCT 60
Qy 61 GCTAATTTCTGTTGGAACTGAACTAACACACCGCGATGAAGTTGATGATCTAGGAAC 120
Db 61 GCTAATTTCTGTTGGAACTGAACTAACACACCGCGATGAAGTTGATGATCTAGGAAC 120
Qy 121 CCTCAAAATCTGTTAATCTGACAAAACCTTTTATTAATATATGCTGCTTCGTT 180
Db 121 CCTCAAAATCTGTTAATCTGACAAAACCTTTTATTAATATATGCTGCTTCGTT 180
Qy 181 CCTGGTCTAGTACGTGTACACCTTGTCCATAAAAAAGATGCTGCTTAACCAAT 240
Db 181 CCTGGTCTAGTACGTGTACACCTTGTCCATAAAAAAGATGCTGCTTAACCAAT 240

Db 181 CCTGGTCTAGTACGTGTACACCTTGTCCATAAAAAAGATGCTGCTTAACCAAT 240
Qy 241 CCACCTGCTACTGCTAAATTTAGTACATAATGTAACGTTAAATGCCCTGCTGGTACCGCA 300
Db 241 CCACCTGCTACTGCTAAATTTAGTACATAATGTAACGTTAAATGCCCTGCTGGTACCGCA 300
Qy 301 ATTGCAGGTGGAGCAACAGATTATGCAGCAATATATGCAGCAATATGTTAATTTAGTAAT 360
Db 301 ATTGCAGGTGGAGCAACAGATTATGCAGCAATATATGCAGCAATATGTTAATTTAGTAAT 360
Qy 361 AATTTTATATGAAATGCTCCAAATTTTAAATGCAGGTGCTAGTACATGACACAGCTTGT 420
Db 361 AATTTTATATGAAATGCTCCAAATTTTAAATGCAGGTGCTAGTACATGACACAGCTTGT 420
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Db 421 CCGGTAACAGAGTTGGTGGTGCATTGACTGCTGTAATGCCGCTACCATAGTCGCATAA 480
Qy 481 TGTAACTGCGCATGCTCCTACTGGTACTGCACTTGTATGATGAGTAACCTGCT 540
Db 481 TGTAACTGCGCATGCTCCTACTGGTACTGCACTTGTATGATGAGTAACCTGCT 540
Qy 541 AGATCAATTCACAGAAATGTTAAATGTAAGTAACTTAACTTTTACTATAATGCTAATAGTGT 600
Db 541 AGATCAATTCACAGAAATGTTAAATGTAAGTAACTTAACTTTTACTATAATGCTAATAGTGT 600
Qy 601 AATACTCCTTTCAATCCAGGTAAAGTTAATGCACACCTTGTCCGCGCAATTAACCTGCT 660
Db 601 AATACTCCTTTCAATCCAGGTAAAGTTAATGCACACCTTGTCCGCGCAATTAACCTGCT 660
Qy 661 AATGTTGCTTAAGCTACTTTAGTAAATGATGCTACATAACCCGCAATATGTAAGTTGCA 720
Db 661 AATGTTGCTTAAGCTACTTTAGTAAATGATGCTACATAACCCGCAATATGTAAGTTGCA 720
Qy 721 TGCCCTGATGCTACTATAAGTGCCTGGAGTAAATTAATTTGGGTAGCACAAAACACTGAA 780
Db 721 TGCCCTGATGCTACTATAAGTGCCTGGAGTAAATTAATTTGGGTAGCACAAAACACTGAA 780
Qy 781 TGTACTAATTTGCTCCTCACTTTTACAATAATTAATGCTCCTAATTTCAATCCAGGTAAT 840
Db 781 TGTACTAATTTGCTCCTCACTTTTACAATAATTAATGCTCCTAATTTCAATCCAGGTAAT 840
Qy 841 AGTACATGCTTACCTCCAGCAAAATAAGATTAATGCTGCTAGAGCCACTGAGGTGGT 900
Db 841 AGTACATGCTTACCTCCAGCAAAATAAGATTAATGCTGCTAGAGCCACTGAGGTGGT 900
Qy 901 GCGCTACTTTAGCCAAATAATGTAATTTGCAATGCTGCTGATGCTGCAATTTGCTAGT 960
Db 901 GCGCTACTTTAGCCAAATAATGTAATTTGCAATGCTGCTGATGCTGCAATTTGCTAGT 960
Qy 961 GGAGCACTAATTTATGTAATTTATTAACAGAAATGCTAATTTGCTGCTGCTAATTTTAT 1020
Db 961 GGAGCACTAATTTATGTAATTTATTAACAGAAATGCTAATTTGCTGCTGCTAATTTTAT 1020
Qy 1021 TTTGATGGTAAATTTCTAGCAGGAAGTAGTAGATGCAAAAGCATGTCAGCAAAATAA 1080
Db 1021 TTTGATGGTAAATTTCTAGCAGGAAGTAGTAGATGCAAAAGCATGTCAGCAAAATAA 1080
Qy 1081 GTTTAAGGCCCTGTAGCAACTGCAGGTGGTACTGCTACTTTAATTTCAATTTGCTGCTT 1140
Db 1081 GTTTAAGGCCCTGTAGCAACTGCAGGTGGTACTGCTACTTTAATTTCAATTTGCTGCTT 1140
Qy 1141 GAATGCCCTGCTGCTACTGCTACTCAGGATGGAACACATCTACTTTAATTAAGCAGCA 1200
Db 1141 GAATGCCCTGCTGCTACTGCTACTCAGGATGGAACACATCTACTTTAATTAAGCAGCA 1200
Qy 1201 TCTGAATGTGTTAAATGCTGCTGCAACTTTTATCTACAAAATAAACTGATTTGGGTAGCA 1260
Db 1201 TCTGAATGTGTTAAATGCTGCTGCAACTTTTATCTACAAAATAAACTGATTTGGGTAGCA 1260
Qy 1261 GGTATTGATACATGCTACTAGTTGTAATAAAAAAATTAACCTTCTGGCGCTGAAGCTAATTTA 1320
Db 1261 GGTATTGATACATGCTACTAGTTGTAATAAAAAAATTAACCTTCTGGCGCTGAAGCTAATTTA 1320

QY 1321 CCGAATCGCTAAATAATATATATATGTCATTTCGCTAAATTTTATCAATTCCTTA 1380
Db 1321 CCTGAATCGCTAAATAATATATATATGTCATTTCGCTAAATTTTATCAATTCCTTA 1380
QY 1381 TTATTGATTCCTTATATTATTA 1404
Db 1381 TTATTGATTCCTTATATTATTA 1404
RESULT 3
AA52136
ID AA52136 standard; DNA; 1404 BP.
XX
AC AA52136;
XX
DT 04-DEC-2000 (first entry)
DE 55 kDa i-antigen gene.
XX
KW BTU1; beta-tubulin; protein expression system; negative selection;
KW paclitaxel sensitivity; cell surface; antigen; protozoa; ciliate;
KW live vaccine; Ichthyophthius multifiliis; immobilization-antigen;
KW i-antigen; freshwater; fish; protozoacide; ds.
XX
OS Ichthyophthius multifiliis.
XX
FH Key Location/Qualifiers
FT CDS 1..1404
FT FT /*tag= a
FT FT /codon= (seq:"TAA", aa:Gln)
FT FT /product= 55_kDa_i-antigen
FT FT /partial
XX
PN W0200046381-A1.
XX
XX 10-AUG-2000.
XX
XX 04-FEB-2000; 2000WO-US02966.
XX
XX 04-FEB-1999; 99US-0118634.
XX 02-MAR-1999; 99US-0122372.
XX 17-MAR-1999; 99US-0124905.
XX 27-APR-1999; 99US-0131121.
XX
XX (UYGE-) UNIV GEORGIA RES FOUND INC.
XX (GAER/) GAERTIG J.
XX (DICK/) DICKERSON H W.
XX (CLAR/) CLARK T G.
XX
XX Gaertig J, Dickerson HW, Clark TG;
XX
XX WPI: 2000-514962/46.
XX P-PSDB: AAY97117.
XX
XX Recombinant expression systems for expressing heterologous nucleic
XX acids and producing recombinant protein, comprises nonpathogenic
XX protozoa such as Tetrahymena resistant to paclitaxel
XX
XX Disclosure; Fig 3B; 83pp; English.
XX
XX Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and
XX BTU2), which encode identical beta-tubulin proteins. Either of these two
XX genes (but not both at once) can be disrupted without a detectable change
XX in the cell phenotype. A K350L substitution in the BTU1 beta-tubulin
XX protein confers increased resistance to microtubule-depolymerizing drugs
XX and increased sensitivity to paclitaxel, a microtubule-stabilizing drug.
XX Cells carrying the BTU1-Ik350M allele can be transformed to paclitaxel
XX resistance by gene replacement of BTU1-Ik350M with a wild-type BTU1 gene
XX fragment, eliminating the need to incorporate a means for positive
XX selection. Where the host organism is not a T. thermophila mutant
XX containing the BTU1-Ik350M allele, BTU1::neol construct, which
XX substitutes the coding region of the neol gene (conferring resistance to

CC parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts
CC and for positive selection. Heterologous nucleic acids (especially
CC encoding antigenic polypeptides) can be inserted into a BTU gene for
CC successful cell-surface expression that is maintained by way of negative
CC selection. Preferred expression vectors disrupt the BTU1-Ik350M gene by
CC homologous recombination-mediated insertion of a heterologous nucleic
CC acid, thereby restoring resistance to paclitaxel in the resulting
CC transgenic host. Transgenic ciliated protozoa are useful as live vaccines
CC for stimulating an immune response in a vertebrate. The transgenic
CC protozoan host cells are also useful for producing polyclonal antibodies
CC (claimed). In particular, Tetrahymena expressing Ichthyophthius
CC multifiliis immobilization-antigen (i-antigen) protein on their surface
CC are effective vehicles for vaccination of freshwater fish against
CC infection by I. multifiliis.

XX
SQ Sequence 1404 BP; 447 A; 241 C; 256 G; 460 T; 0 other;

Query Match 99.9%; Score 1402.4; DB 21; Length 1404;

Best Local Similarity 99.9%; Pred. No. 3.4e-301;

Matches 1403; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 ATGAAAAATAATTTTAGTAATATTGATTATTTTCATTATTTATCAATTTAAATTAATCT 60

Db 1 ATGAAAAATAATTTTAGTAATATTGATTATTTTCATTATTTATCAATTTAAATTAATCT 60

QY 61 GCTAATTGCTGTTGGAACTGAAACTAACACAGCCGGATAGTTGATGATCTAGGAAC 120

Db 61 GCTAATTGCTGTTGGAACTGAAACTAACACAGCCGGATAGTTGATGATCTAGGAAC 120

QY 121 CTGCAAAATGTTAAATTTGTAGAAAACTTTTATTATAATATGCTGCTTCGTTGTT 180

Db 121 CTGCAAAATGTTAAATTTGTAGAAAACTTTTATTATAATATGCTGCTTCGTTGTT 180

QY 181 CTTGTTGCTAGTACGTTACACCTTTTCCATAAAAAAAGATGCTGCTTAACCAAT 240

Db 181 CTTGTTGCTAGTACGTTACACCTTTTCCATAAAAAAAGATGCTGCTTAACCAAT 240

QY 241 CCACCTGCTACTGCTAAATTTAGTACATAATGTAACGTTAAATGCCCTGGTACC 300

Db 241 CCACCTGCTACTGCTAAATTTAGTACATAATGTAACGTTAAATGCCCTGGTACC 300

QY 301 ATTGCAGTGGAGCAACAGATTATGCAGCAATAATCAGCAATGTTAATTTGAGAAT 360

Db 301 ATTGCAGTGGAGCAACAGATTATGCAGCAATAATCAGCAATGTTAATTTGAGAAT 360

QY 361 AATTTTATAATGAAAAATGCTCCAAATTTTAAATGAGGCTAGTACATGACACGCTT 420

Db 361 AATTTTATAATGAAAAATGCTCCAAATTTTAAATGAGGCTAGTACATGACACGCTT 420

QY 421 CCGGTAACACAGAGTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 480

Db 421 CCGGTAACACAGAGTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 480

QY 481 TGTAACTGCTCATGCTCTACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 540

Db 481 TGTAACTGCTCATGCTCTACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 540

QY 541 AGATCAATTCACAAATGCTGTTAAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 600

Db 541 AGATCAATTCACAAATGCTGTTAAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 600

QY 601 AATGCTCCTTTCAATCCAGTAAAGTTAATGCACACCTTTGTCGGGCAATTAACCTGCT 660

Db 601 AATGCTCCTTTCAATCCAGTAAAGTTAATGCACACCTTTGTCGGGCAATTAACCTGCT 660

QY 661 AATGTTGCTTAAAGTACTTCTAGTAAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 720

Db 661 AATGTTGCTTAAAGTACTTCTAGTAAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 720

QY 721 TGCCTCATGCTACTAAAGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 780

Db 721 TGCCTCATGCTACTAAAGTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 780

Qy 121 CCTCAAAATGTGTTAAATTTAGAAAAAATTTTATTATATATATGCTGCTTTCGTT 180
Db 121 CCTGCTAACGTGTGAACTGTGACAGAAACTTCTACTACAAACACGCTGTGCTTTCGTTG 180
Qy 181 CCTGGTGTAGTGTGTACACGTGTGCTTCCATAAAAAAGATGCTGTGTCTTAAACCAAAAT 240
Db 181 CCTGGAGCTTCTACCTGTACCCCTTGTCTCAGAGAAGACGCTGGAGCTCAGCCTAAC 240
Qy 241 CCACCTGCTACTGCTAATTTAGTACACATAATGTAAAGCTTAAATGCCCCCTGCTGACGCA 300
Db 241 CCTCTGCTACCGCTAACCTGGTGACCCAGTGAAGCTGAAAGTGTCTGCTGGAACCGCT 300
Qy 301 ATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACGAATGTGTAAATGTAGAATT 360
Db 301 ATCGCTGGAGGAGTACCGACTACCGTGTATCATCACCGAGTGTGTAACCTGTCGCATC 360
Qy 361 AATTTTATATGAATGAAATGCTCCAAATTTTAAATGCAGGTGCTAGTACATGCACAGCTTGT 420
Db 361 AACTTCTACACGAGAACGCTCCTAACTTCAACGCTGGAGCTTCTACCTGTACCGCTTGT 420
Qy 421 CCGGTAACACAGATTGCTGTGCATTTGACTGCTGGTAATGCCGCTACCATAGTCGCATAA 480
Db 421 CCTGTGAACCGCTGGAGAGCTGTGACCGCTGGAAACGCTGCTACCATCGTGGCTCAG 480
Qy 481 TGTAACTGCGCATGCTCTACTGTGCTACTGCATTTGATGGATGAATACTACTGATTATGTT 540
Db 481 TGTAACTGGCTTGTCTCTACCGAACCGCTCTGGAGACGGAGTGACCAACCGACTACGTG 540
Qy 541 AGATCAATTCACAGAATGTGTTAAATGTAGACTTAACCTTTTACTATATGTTAAATGTT 600
Db 541 CGTCTTTTACCAGTGTGTGAAGTGTGCGCTGAACTTCTACTACAAACGAAACACGGA 600
Qy 601 AATACTCTTTTCAATCCAGGTAAAGTAAATGCACACCTTGTCCGCAATTTAAACCTGCT 660
Db 601 AACACCCCTTTCAACCCCTGAAAGTCTCAGTGTACCCCTTGTCTGCTATCAAGCTGCT 660
Qy 661 AATGTTGCTTAACTACTTACTAGTATGATGCTACAAATACCGCATAAATGTAACGTGCA 720
Db 661 AAGTGGCTCAGGCTACCCCTGGGAACGACGCTACCATCAGCGCTCAGTGTAACTGGCT 720
Qy 721 TGCCCTGATGGTACTATAAGTGTGCTGGAGTAAATAATTTGGGTAGCACAACACTGAA 780
Db 721 TGTCTCTGACGGAACCATCTCTGCTGCTGGAGTGAACACTGGGTGGCTCAGAACCCGAG 780
Qy 781 TGTACTAATTTGCTCCTAACTTTTACAAATAATATGCTCCTTAATTTCAATCCAGGTAAT 840
Db 781 TGTACCAACTGTGCTCCTAACTTCTACAAACAACGCTCCTTAATTTCAACCTGGAAC 840
Qy 841 AGTACATGCTACCTTGCCACCAATTAAGATTATGCTGTAAGCCACTGCAGGTGGT 900
Db 841 TCTACTGTCTGCTGCTGCTGCTTGAACAGGACTACGAGCTGAGGCTACCGCTGGAGGA 900
Qy 901 GCCGTACTTTTACGCAAAATATGTAATATGTCATGCCCTGATGCTGCTGCTGCTGCTAGT 960
Db 901 GCTGTACCTTACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 960
Qy 961 GGAGCAACTAATTTATATATATAAACAAGATGTCTTAAATTTGCTGCTGCTGCTGCTGCT 1020
Db 961 GGAGCTACCAACTAGCTGATCCTGACAGACCGAGTGTCTGAACTGTGCTGCTGCTGCTGCT 1020
Qy 1021 TTTGATGTAATATTTCTAGCAGGAAGTAGTAGATGCAAGCATGTCCAGCAAAATAA 1080
Db 1021 TTCAGCGGAACAACCTTCCAGGCTGGATCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1080
Qy 1081 GTTTAAGCGCTGTAGCAACTGCAGGTGGTACTGCTGCTTAAATTTGATATGTCGCTT 1140
Db 1081 GTGACGGGAGCTGTGGCTTACCGCTGGAGAACCGCTACCCCTGATGCTGCTGCTGCTGCT 1140
Qy 1141 GAATGCCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1200
Db 1141 GAGTGTCTGCTGGAACCGTGTGACCGACGGAACCACTCTACCTACAAACGAGCTGCT 1200
Qy 1201 TCTGAATGTGTTAAATGTGCTGCCAACTTTTATACTACAAATAAATGATGGGTAGCA 1260

Db 1201 TCTGAGTGTGTGAAGTGTGCTGCTAACTTCTACACCACCAAGCAGACCGAGCTGGTGGCT 1260
Qy 1261 GGTATTGATACATGACTAGTTGCTTAATAAAAAATTAACATTTCTGGCGCTGAAGCTAATTTA 1320
Db 1261 GGAATCGACACCTGTACCTCTTGTGAACAAGAAGCTGACCTCTGGAGCTGAGGCTAACCTG 1320
Qy 1321 CCTGAATCTGCTAAAAAATAATATAATGATGTTTCGGCTAAATTTTATCAATTTTCCCTTA 1380
Db 1321 CCTGAGTCTGCTAAGAAGAACAATCCAGTGTGACTTCGCTAACTTCTGCTATCTCTCTG 1380
Qy 1381 TTATTGATTTCTTATTA 1397
Db 1381 CTGCTGATCTCTTACTA 1397

RESULT 6

AAA97065

ID AAA97065 standard; DNA; 1404 BP.

AC AAA97065;

XX 18-DEC-2000 (first entry)

XX Synthetic 55KD i-antigen gene sequence.

XX Immobilisation antigen; i-antigen; Ichthyophthiriasis; vaccine; ds;

KW white spot disease; freshwater fish; Immune response; Infection control.

XX Ichthyophthirius multifiliis.

OS Synthetic.

XX WO200046373-A1.

PD 10-AUG-2000.

XX 04-FEB-2000; 2000WO-US02962.

XX 04-FEB-1999; 99US-0118634.

PR 02-MAR-1999; 99US-0122372.

PR 17-MAR-1999; 99US-0124905.

PR 27-APR-1999; 99US-0131121.

XX (UIGE-) UNIV GEORGIA RES FOUND INC.

PA (CORR) CORNELL RES FOUND INC.

PA (CLAR) CLARK T G.

PA (DICK) DICKERSON H W.

PA (LINT) LIN T.

PI Clark TG, Dickerson HW, Lin T;

XX WPI; 2000-0506071/45.

Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish .

Example 5; Figure 13; 144pp; English.

This invention relates to novel i-antigen polypeptide sequences. I-antigens or immobilisation antigens are common to a variety of hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in Ichthyophthirius multifiliis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kD. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis serotypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an

CC antigenic i-antigen polypeptide obtained is also useful for treating or
CC preventing I. multifiliis infection in fish. Sequences AAA97036-A97042,
CC and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene
CC fragments identified in the invention. Sequences AAA97043-A97064
CC (excluding AAA97060) and AAA97071-A97088 represent primers used in the
CC isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and
CC AAB25893-B25906 represent i-antigen protein and peptide sequences.

XX
SQ Sequence 1404 BP; 317 A; 418 C; 339 G; 330 T; 0 other;

Query Match 55.6%; Score 781; DB 21; Length 1404;
Best Local Similarity 72.4%; Pred. No. 1.6e-163;
Matches 1012; Conservative 0; Mismatches 385; Indels 0; Gaps 0;

QY 1 ATGAAATAATATTTAGTAATATTTGATATTTTCAATTTATCAATTAATTAATCT 60
DB 1 ATGAAGAACAACATCCGGTGATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 60
QY 61 GCTAATTGCTCTTGGAACTGAACTAAACACACGCGGATAAGTTGATGATCTAGGAAT 120
DB 61 GCTAATGCTCTTGGAACTGAACTAAACACACGCGTGGACAGTGGACACCTGGGAACC 120
QY 121 CCTGCAATTTGTTAATTTAGAAAACTTTTATTAATTAATTAATGCTGCTGCTTCGTT 180
DB 121 CCTGCAATTTGTTAATTTAGAAAACTTTTATTAATTAATTAATGCTGCTGCTTCGTT 180
QY 181 CCTGGTCTAGTACGCTGACACCTTGTCCATAAAAAAGATGCTGGTGCCTTAACCAAT 240
DB 181 CCTGGTCTAGTACGCTGACACCTTGTCCATAAAAAAGATGCTGGTGCCTTAACCAAT 240
QY 241 CCACCTGCTACTGCTAATTTAGTACATATGTAATGTAATGTAATGTAATGTAATGTAAT 300
DB 241 CCACCTGCTACTGCTAATTTAGTACATATGTAATGTAATGTAATGTAATGTAATGTAAT 300
QY 301 ATTCAGGTGGGACACAGATTTATGACGCAATTAATCACAGAAATGCTTTAATTTGAAAT 360
DB 301 ATTCAGGTGGGACACAGATTTATGACGCAATTAATCACAGAAATGCTTTAATTTGAAAT 360
QY 361 AATTTTATATGAAATGCTCCAAATTTTAAATGCAAGTCTAGTACATACACAGCTTGT 420
DB 361 AATTTTATATGAAATGCTCCAAATTTTAAATGCAAGTCTAGTACATACACAGCTTGT 420
QY 421 CCGGTAACAGAGATGTTGGTGCATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 480
DB 421 CCGGTAACAGAGATGTTGGTGCATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 480
QY 481 TGTACGTCGATGCTCTACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 540
DB 481 TGTACGTCGATGCTCTACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 540
QY 541 AGATCATTACAGAGATGTTGTTAATGTAAGTAACTTAACTTTTAACTTAACTTAACTTAA 600
DB 541 AGATCATTACAGAGATGTTGTTAATGTAAGTAACTTAACTTTTAACTTAACTTAACTTAA 600
QY 601 AATACCTCTTCAATCCAGGTAAAGTTAATGCAACCTTGTCCGCAATTAACCTGCT 660
DB 601 AATACCTCTTCAATCCAGGTAAAGTTAATGCAACCTTGTCCGCAATTAACCTGCT 660
QY 661 AATGCTGCTTAAAGTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACT 720
DB 661 AATGCTGCTTAAAGTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACT 720
QY 721 TGCCCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 780
DB 721 TGCCCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 780
QY 781 TGTACTAATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 840
DB 781 TGTACTAATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 840
QY 841 AGTACATGCTACCTTGGCCAGCAATTAAGATTTATGCTGCTGCTGCTGCTGCTGCTGCTG 900
DB 841 AGTACATGCTACCTTGGCCAGCAATTAAGATTTATGCTGCTGCTGCTGCTGCTGCTGCTG 900

QY 901 GCCCTACTTTAGCCAAATTAATGTAATTAATGCAATGCCCTGATGGTACTGCAATTTGCTAGT 960
DB 901 GCTGCTACCTTGGCTAAGCAGTGTAAATGCAATGCCCTGATGGTACTGCAATTTGCTAGT 960
QY 961 GGACCACTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 1020
DB 961 GGACCACTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 1020
QY 1021 TTTGATGTAATAATTTCTAGGAGGAGTATAGTATGCAAGCATGTCAGCAAGATAA 1080
DB 1021 TTTGATGTAATAATTTCTAGGAGGAGTATAGTATGCAAGCATGTCAGCAAGATAA 1080
QY 1081 GTTTAAGCGCTGTAGCAACTGCAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1140
DB 1081 GTTTAAGCGCTGTAGCAACTGCAAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1140
QY 1141 GAATGCCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1200
DB 1141 GAATGCCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1200
QY 1201 TCTGAATGTTAAATGCTGCCAATTTTATATACAAAATTAATGATGTTGGTAGCA 1260
DB 1201 TCTGAATGTTAAATGCTGCCAATTTTATATACAAAATTAATGATGTTGGTAGCA 1260
QY 1261 GGTATTGATACATGCTAGTGTGTAATTAATTAATTAATTAATTAATTAATTAATTA 1320
DB 1261 GGTATTGATACATGCTAGTGTGTAATTAATTAATTAATTAATTAATTAATTAATTA 1320
QY 1321 CCTGAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1380
DB 1321 CCTGAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1380
QY 1381 TTATTGATTTCTTATTA 1397
DB 1381 TTATTGATTTCTTATTA 1397
RESULT 7
AAA97036 standard; DNA; 1326 BP.
XX AAA97036;
XX 18-DEC-2000 (first entry)
XX 48kd i-antigen nucleotide sequence.
XX Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; ds;
KW white spot disease; freshwater fish; immune response; infection control.
XX Ichthyophthirius multifiliis.
XX WO200046373-A1.
XX 10-AUG-2000.
XX 04-FEB-2000; 2000WO-US02962.
XX 04-FEB-1999; 99US-0118634.
XX 02-MAR-1999; 99US-0122372.
XX 17-MAR-1999; 99US-0124905.
XX 27-APR-1999; 99US-0131121.
XX (UYGE-) UNIV GEORGIA RES FOUND INC.
PA (CORR) CORNELL RES FOUND INC.
PA (CLAR) CLARK T G.
PA (DICK) DICKERSON H W.
XX (LINT) LIN T.
XX Clark TG, Dickerson HW, Lin T;
XX WPI; 2000-506071/45.
DR

PA (LINT/) LIN T.
XX Clark TG, Dickerson HW, Lin T;
XX WPI: 2000-506071/45.
DR
XX Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius
PT multifiliis, useful for prophylaxis and treatment of Ichthyophthirius
PT infection in fish -
PT
XX Disclosure: Figure 1; 144pp; English.
PS
XX This invention relates to novel i-antigen polypeptide sequences.
CC I-antigens or immobilisation antigens are common to a variety of
CC hymenostomatid ciliates and their expression varies in response to
CC environmental stimuli. This invention relates to i-antigens in
CC Ichthyophthirius multifiliis, a protozoan which is an obligate parasite
CC of freshwater fish causing ichthyophthiriasis or white spot disease. The
CC invention includes two polypeptide and polynucleotide sequences for two
CC i-antigens, of 48 and 55 kD. Also included in the invention are
CC antibodies capable of binding to the nucleotide sequences and a method
CC for identifying I. multifiliis serotypes using the nucleotide sequences.
CC A composition (containing the i-antigen nucleotide) capable of eliciting
CC an immune response in fish is useful for prophylaxis, treatment or for
CC controlling I. multifiliis infection in fish. Polynucleotide or protein
CC vaccines comprising a portion of the amplified product encoding an
CC antigenic i-antigen polypeptide obtained is also useful for treating or
CC preventing I. multifiliis infection in fish. Sequences AAA97036-A97042,
CC and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene
CC fragments identified in the invention. Sequences AAA97043-A97064
CC (excluding AAA97060) and AAA97071-A97088 represent primers used in the
CC isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and
CC AAB25893-B25906 represent i-antigen protein and peptide sequences.
XX
SQ Sequence 2486 BP; 896 A; 310 C; 321 G; 959 T; 0 other;
Query Match 18.0%; Score 252.6; DB 21; Length 2486;
Best Local Similarity 56.7%; Pred. No. 2.2e-46;
Matches 660; Conservative 0; Mismatches 394; Indels 111; Gaps 6;
QY 344 GTGTTAATCTAGAAATTTTATATGAAGAAATGCTCCAAATTTTATGCAGGTGCTA 403
DB 599 GTGCTGCTTAAGGAGAGCTAATGTTGTAATTAACCTTTCGACCAATATATGCTGAG 658
QY 404 GTACATGCAGCTGTGCGGTAAACAGAGTTGGTGGTGCATGTGCTGTTAATGCCG 463
DB 559 GTATATGTGTACCATGCCAATAAACAAGAGTAGGCTCTGTACCAATGCAGGTGACTTAG 718
QY 464 CTACCATAGTCGGATATGTAAGCGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 523
DB 719 CTACTTTAGCCACATAATGCAGTACTTAATGTCTACTGCGCACTGCACCTGATGATGG 778
QY 524 TAACCTAGTATGTTAGATCAATTCACAGAGTGTCTTAAATGTAGACTTAACCTTTTACT 583
DB 779 TGACAGATGTTTGTATAGATCAGCCGCAATATGCTTAAATGCAACCTTAACCTTTACT 838
QY 584 ATATGGTATAATAGTAAATCTCCTTTCAATCCAGTAAAAAGTTAATGCACACCTTGTC 643
DB 839 ATATGGTGGTCTCCCTTAAGGTGAAGCTCCTGGCGTTTAAAGTTTGTGCTGCTGCTG 898
QY 644 CGGCAATTAACCTGCT-----AATG 664
DB 899 CCGCTGCAGGTGTTGCTGCGCTTACTAGTTAATGTGTACCTGCCAATTAACAAACG 958
QY 665 TTGCTTAAGCTACTTTAGTAAATGATGTACATAACCGCATATGCTAAGCTTGCATGCC 724
DB 959 ATTCTCTGCCACTTGCAGGTGCCCTAACTAATTTAGCCACATAATGTAGCAATTAATGTC 1018
QY 725 CTGATGCTATTAAGTGTGCTGGAGT----AAATAATGGGTAGCACAACACACTGAT 781
DB 1019 CTACTGGCACTGTACTTGATGATGAGTGACACTTGTGTTTAAATATACATGAGCAAT 1078
QY 782 GTACTAATGTGCTCCCAACTTTTACATAATAATATGCTCCTAAT-----826

DB 1079 GTGTTAATGCAGACCTAACTTTTACTATAATGTTGTTCTCTCTTAAGTGAAGCTCCTG 1138
QY 827 -----TCAATCCAGGTAATAGTACAT 847
DB 1139 GCGTTTAAGTTTTTGTGCTGCTGCCGCTGCCGCTGAGGTGTTGCTGCCGTTACTAGTAAT 1198
QY 848 GCCTACCTTGCAGCAAAATAAGATTATGTTGCTGAAGCCACTGCAGGTGGTGGCGCTA 907
DB 1199 GTGPACCTTGGCAATAAACAACAAACGATCTCTCT---GCCACTGCAGGTGCCAAGCTA 1255
QY 908 CTTTAGCCAAATAATGTAAATATGTCATGCCCTGATGTACTGCAATTTGCTAGTGGAGCA 967
DB 1256 ATTTAGCCACATAATGCAGTACTTAATGTCCACTGCAATTTCAAGCGGAGTGA 1315
QY 968 CTAATATGTAATATTAACACAGAAATGCTFAAATTTGCTGCTAACTTTTATTTTATG 1027
DB 1316 CACTTGTGTTTGTAGTAATTCATCCACATAATGTTCTTAATGCATGCTAATTAATCT 1375
QY 1028 GTAATAATTTCTAGCGAAGTAGTAGATCAAGCAAGATGTCAGCAAAATAAGTTTAA 1087
DB 1376 ATGGTAATTTTGAAGCAGGTAAGCTTAATGTTAAAGTGTCCAGTAAGTAAACT---A 1432
QY 1088 GCGCTGTAGCAACTGCAGGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1147
DB 1433 CTCCAGCACATGCTCCAGGTAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1492
QY 1148 CTGCTGGTACTGCTACCGATGGAACAACATCTACTTATAATAAGCAGCATCTGAAT 1207
DB 1493 CTGCTGGTACACTGCTGATGATGGAACATCAACTTAATTTTGTAGCTTCCGCAACTGA 1552
QY 1208 GTGTTAAATGCTGCTGCAACTTTTATATACAAAATAAATGATGTTGGTACGAGTATG 1267
DB 1553 GTACTAATGTTCTGCTGGCTTTTGTGCATCAAAACAACTGGTTTACACGAGTACTG 1612
QY 1268 ATACATGCTACTAGTTGTAATAAAAAATTAACCTTCTGGCGCTGAAGCTAATTTACCTGA 1327
DB 1613 ATACATGCTACTGAATGCTACTAAAAATTAACCTTCTGGTGCACAGCTAAAGTATATGCTG 1672
QY 1328 CTGCTAAAAAATAATATAATG-----TGATTTGCTAAATTTTATCAATTTCT 1378
DB 1673 AGCTACTCAAAAGATATAATGCGCTCCACTACTTTCGCTAAATTTTATCGATTTCCT 1732
QY 1379 TATATTGATTTCTTATTTATTTATT 1403
DB 1733 TATATTATTTCTTCTATTTATT 1757
RESULT 9
AAA52135
ID AAA52135 standard; DNA; 1326 BP.
XX AAA52135;
XX XX
DT 04-DEC-2000 (first entry)
XX
DE 48 kDa i-antigen gene.
XX
KW BTU1; beta-tubulin; protein expression system; negative selection;
KW pacilitaxel sensitivity; cell surface; antigen; protozoa; ciliate;
KW live vaccine; Ichthyophthirius multifiliis; immobilization-antigen;
KW i-antigen; freshwater; fish; protozoacide; ds.
XX
OS Ichthyophthirius multifiliis.
XX
FH Key Location/Qualifiers
FT CDS 1..1326
FT /*tag= a
FT /transl_except= "pos:82...84, aa:Gln"
FT /codon= (seq:"TAA", aa:Gln)
FT /product= 48_kDa_i-antigen
FT /partial
XX


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XX (UYGE-) UNIV GEORGIA RES FOUND INC.
PA (CORR ) CORNELL RES FOUND INC.
PA (CLARK/) CLARK T G.
PA (DICK/) DICKERSON H W.
PA (LINT/) LIN T.
XX Clark TG, Dickerson HW, Lin T;
PI WPI; 2000-506071/45.
XX
XX Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius
PT multifiliis, useful for prophylaxis and treatment of Ichthyophthirius
PT infection in fish -
XX
XX Disclosure; Figure 12; 144pp; English.
XX
XX This invention relates to novel i-antigen polypeptide sequences.
CC I-antigens or immobilisation antigens ar common to a variety of
CC hymenostomatid ciliates and their expression varies in response to
CC environmental stimuli. This invention relates to i-antigens in
CC Ichthyophthirius multifiliis, a protozoan which is an obligate parasite
CC of freshwater fish causing Ichthyophthiriasis or white spot disease. The
CC invention includes two polypeptide and polynucleotide sequences for two
CC i-antigens, of 48 and 55 kD. Also included in the invention are
CC antibodies capable of binding to the nucleotide sequences and a method
CC for identifying I. multifiliis serotypes using the nucleotide sequences.
CC A composition (containing the i-antigen nucleotide) capable of eliciting
CC an immune response in fish is useful for prophylaxis, treatment or for
CC controlling I. multifiliis infection in fish. Polynucleotide or protein
CC vaccines comprising a portion of the amplified product encoding an
CC antigenic i-antigen polypeptide obtained in fish. Sequences AAA97036-A97042,
CC and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene
CC fragments identified in the invention. Sequences AAA97043-A97064
CC (excluding AAA97060) and AAA97071-A97088 represent primers used in the
CC isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and
CC AAB25893-B25906 represent i-antigen protein and peptide sequences.
XX
XX Sequence 100 BP; 16 A; 35 C; 24 G; 25 T; 0 other;
SQ
Query Match 4.5%; Score 62.8; DB 21; Length 100;
Best Local Similarity 77.6%; Pred. No. 0.0001;
Matches 76; Conservative 0; Mismatches 22; Indels 0; Gaps 0;
QY 166 GCGCTGCTTCGTTCTGCTGCTAGTACGTGTACACCTTGTCATATAAAAAAGATGCT 225
DB 2 GCTGCTGCTTTCTGCTGCTGAGCTTCTACCTGTACCCCTTGCTCAGAGAAGGACGCT 61
QY 226 GGTGCTTAACCAATCCACCTGCTACTGCTAATTTAGT 263
DB 62 GGAGCTCAGCGCTAACCGCTCTGCTACCGCTAACCTGGT 99
RESULT 15
AAA97080/c
ID AAA97080 standard; DNA; 100 BP.
XX
XX AAA97080;
AC
XX
XX 18-DEC-2000 (first entry)
DT
XX
XX G5 synthetic gene synthesis primer 3210.
DE
XX
XX Immobilisation antigen; i-antigen; Ichthyophthiriasis; vaccine;
KW white spot disease; freshwater fish; immune response; infection control;
KW PCR primer; ss.
XX
XX Synthetic.
OS
XX
XX WO200046373-A1.
PN
XX
XX 10-AUG-2000.
PD

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XX 04-FEB-2000; 2000WO-US02962.
PF
XX
XX 04-FEB-1999; 99US-0118634.
PR
XX 02-MAR-1999; 99US-0122372.
PR
XX 17-MAR-1999; 99US-0124905.
PR
XX 27-APR-1999; 99US-0131121.
XX
XX (UYGE-) UNIV GEORGIA RES FOUND INC.
PA (CORR ) CORNELL RES FOUND INC.
PA (CLARK/) CLARK T G.
PA (DICK/) DICKERSON H W.
PA (LINT/) LIN T.
XX
XX Clark TG, Dickerson HW, Lin T;
PI WPI; 2000-506071/45.
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CC an immune response in fish is useful for prophylaxis, treatment or for
CC controlling I. multifiliis infection in fish. Polynucleotide or protein
CC vaccines comprising a portion of the amplified product encoding an
CC antigenic i-antigen polypeptide obtained in fish. Sequences AAA97036-A97042,
CC and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene
CC fragments identified in the invention. Sequences AAA97043-A97064
CC (excluding AAA97060) and AAA97071-A97088 represent primers used in the
CC isolation of the i-antigen gene sequences. Sequences AAB25859-B25889 and
CC AAB25893-B25906 represent i-antigen protein and peptide sequences.
XX
XX Sequence 100 BP; 22 A; 17 C; 32 G; 29 T; 0 other;
SQ
Query Match 4.5%; Score 62.8; DB 21; Length 100;
Best Local Similarity 77.6%; Pred. No. 0.0001;
Matches 76; Conservative 0; Mismatches 22; Indels 0; Gaps 0;
QY 754 AATAATTGGGTAGCACAAACACTGAATGTACTAATTTGCTCTCTTAACATTAAT 813
DB 99 AACAACTGGGTGGCTCAGAACACCCAGGTGTACCAACTGTCTCTTAACATTAAC 40
QY 814 AATGCTCTCTAATTCATCCAGGTAATAGTACATCCCT 851
DB 39 AACGCTCTCTAATTCACACCTGGAAACTCTACCTGTCT 2
Search completed: February 16, 2003, 17:00:21
Job time : 224.022 secs

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